## 230 O'd PCT/PTO 1 6 JUN 2000

## SEQUENCE LISTING

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Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp Asp Pro Ile Asn Asn Lys 100 105 110

Asp Leu Phe Ala Cys Lys Arg Ile Leu Leu Lys Ala Lys Ala Leu Asp 115 120 125

Tyr Ile Asn Ala Ser Val Ala Gly Gly Ser Asp Phe Leu Asn Asp Tyr 130 135 140

Val Leu Lys Tyr Ser Glu Arg Tyr Glu Thr Arg Arg Arg Val Gln Ser 145 150 155 160

Thr Gly Arg Ala Lys Ala Ala Phe Asp Glu Asp Gly Asn Ile Ser Asn 165 170 175

Val Lys Ser Asp Thr Asp Arg Asp Ala Glu Thr Val Thr Gln Asn Glu
180 185 190

Asp Ala Asp Arg Asn Asn Ser Ser Arg Ile Asn Leu Gln Asp Ile Cys 195 200 205

Lys Asp Leu Asn Gln Glu Gly Asp Asp Gln Met Phe Val Arg Lys Thr 210 215 220

Ser Arg Glu Ile Gln Gly Gln Asn Arg Arg Asn Leu Ile His Glu Asp 225 230 235 240

Ile Ala Lys Ala Ile Lys Gln Leu Trp Asn Asn Asp Lys Gly Ile Lys 245 250 255

Gln Cys Phe Ala Arg Ser Asn Glu Phe Gln Leu Glu Gly Ser Ala Ala 260 265 270

Tyr Tyr Phe Asp Asn Ile Glu Lys Phe Ala Ser Pro Asn Tyr Val Cys 275 280 285 Thr Asp Glu Asp Ile Leu Lys Gly Arg Ile Lys Thr Thr Gly Ile Thr 290 295 300

Glu Thr Glu Phe Asn Ile Gly Ser Ser Lys Phe Lys Val Leu Asp Ala 305 310 315 320

Gly Gly Gln Arg Ser Glu Arg Lys Lys Trp Ile His Cys Phe Glu Gly 325 330 335

Ile Thr Ala Val Leu Phe Val Leu Ala Met Ser Glu Tyr Asp Gln Met 340 345 350

Leu Phe Glu Asp Glu Arg Val Asn Arg Met His Glu Ser Ile Met Leu 355 360 365

Phe Asp Thr Leu Leu Asn Ser Lys Trp Phe Lys Asp Thr Pro Phe Ile 370 375 380

Leu Phe Leu Asn Lys Ile Asp Leu Phe Glu Glu Lys Val Lys Ser Met 385 390 395 400

Pro Ile Arg Lys Tyr Phe Pro Asp Tyr Gln Gly Arg Val Gly Asp Ala 405 410 415

Glu Ala Gly Leu Lys Tyr Phe Glu Lys Ile Phe Leu Ser Leu Asn Lys 420 425 430

Thr Asn Lys Pro Ile Tyr Val Lys Arg Thr Cys Ala Thr Asp Thr Gln 435

Thr Met Lys Phe Val Leu Ser Ala Val Thr Asp Leu Ile Ile Gln Gln 450 455 460

Asn Leu Lys Glu Tyr Asn Leu Val 465 470

<210> 109

<211> 472

<212> PRT

<213> Chimaera sp.

<400> 109

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro 1 5 10 15

Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln 20 25 30

Leu Glu Lys Gln Arg Asp Lys Asn Glu Ile Lys Leu Leu Leu Gly 35 40 45

Ala Gly Glu Ser Gly Lys Ser Thr Val Leu Lys Gln Leu Lys Leu Leu 50 55 60

His Gln Gly Gly Phe Ser His Gln Glu Arg Leu Gln Tyr Ala Gln Val 65 70 75 80

- Ile Trp Ala Asp Ala Ile Gln Ser Met Lys Ile Leu Ile Ile Gln Ala 85 90 95
- Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp Asp Pro Ile Asn Asn Lys
  100 105 110
- Asp Leu Phe Ala Cys Lys Arg Ile Leu Leu Lys Ala Lys Ala Leu Asp 115 120 125
- Tyr Ile Asn Ala Ser Val Ala Gly Gly Ser Asp Phe Leu Asn Asp Tyr 130 135 140
- Val Leu Lys Tyr Ser Glu Arg Tyr Glu Thr Arg Arg Arg Val Gln Ser 145 150 155 160
- Thr Gly Arg Ala Lys Ala Ala Phe Asp Glu Asp Gly Asn Ile Ser Asn 165 170 175
- Val Lys Ser Asp Thr Asp Arg Asp Ala Glu Thr Val Thr Gln Asn Glu 180 185 190
- Asp Ala Asp Arg Asn Asn Ser Ser Arg Ile Asn Leu Gln Asp Ile Cys 195 200 205
- Lys Asp Leu Asn Gln Glu Gly Asp Asp Gln Met Phe Val Arg Lys Thr 210 215 220
- Ser Arg Glu Ile Gln Gly Gln Asn Arg Arg Asn Leu Ile His Glu Asp 225 230 235 240
- Ile Ala Lys Ala Ile Lys Gln Leu Trp Asn Asn Asp Lys Gly Ile Lys 245 250 255
- Gln Cys Phe Ala Arg Ser Asn Glu Phe Gln Leu Glu Gly Ser Ala Ala 260 265 270
- Tyr Tyr Phe Asp Asn Ile Glu Lys Phe Ala Ser Pro Asn Tyr Val Cys 275 280 285
- Thr Asp Glu Asp Ile Leu Lys Gly Arg Ile Lys Thr Thr Gly Ile Thr 290 295 300
- Glu Thr Glu Phe Asn Ile Gly Ser Ser Lys Phe Lys Val Leu Asp Ala 305 310 315 320
- Gly Gly Gln Arg Ser Glu Arg Lys Lys Trp Ile His Cys Phe Glu Gly 325 330 335
- Ile Thr Ala Val Leu Phe Val Leu Ala Met Ser Glu Tyr Asp Gln Met 340 345 350
- Leu Phe Glu Asp Glu Arg Val Asn Arg Met His Glu Ser Ile Met Leu 355 360 365
- Phe Asp Thr Leu Leu Asn Ser Lys Trp Phe Lys Asp Thr Pro Phe Ile 370 375 380
- Leu Phe Leu Asn Lys Ile Asp Leu Phe Glu Glu Lys Val Lys Ser Met

Pro Ile Arg Lys Tyr Phe Pro Asp Tyr Gln Gly Arg Val Gly Asp Ala 405 410 415

Glu Ala Gly Leu Lys Tyr Phe Glu Lys Ile Phe Leu Ser Leu Asn Lys 420 425 430

Thr Asn Lys Pro Ile Tyr Val Lys Arg Thr Cys Ala Thr Asp Thr Gln
435
440
445

Thr Met Lys Phe Val Leu Ser Ala Val Thr Asp Leu Ile Ile Gln Gln 450 455 460

Asn Leu Lys Asp Ile Met Leu Gln 465 470

<210> 110

<211> 472

<212> PRT

<213> Chimaera sp.

<400> 110

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro 1 5 10 15

Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln 20 25 30

Leu Glu Lys Gln Arg Asp Lys Asn Glu Ile Lys Leu Leu Leu Gly 35 40 45

Ala Gly Glu Ser Gly Lys Ser Thr Val Leu Lys Gln Leu Lys Leu Leu 50 55 60

His Gln Gly Gly Phe Ser His Gln Glu Arg Leu Gln Tyr Ala Gln Val 65 70 75 80

Ile Trp Ala Asp Ala Ile Gln Ser Met Lys Ile Leu Ile Ile Gln Ala 85 90 95

Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp Asp Pro Ile Asn Asn Lys 100 105 110

Asp Leu Phe Ala Cys Lys Arg Ile Leu Leu Lys Ala Lys Ala Leu Asp 115 120 125

Tyr Ile Asn Ala Ser Val Ala Gly Gly Ser Asp Phe Leu Asn Asp Tyr 130 135 140

Val Leu Lys Tyr Ser Glu Arg Tyr Glu Thr Arg Arg Arg Val Gln Ser 145 150 155 160

Thr Gly Arg Ala Lys Ala Ala Phe Asp Glu Asp Gly Asn Ile Ser Asn 165 170 175

Val Lys Ser Asp Thr Asp Arg Asp Ala Glu Thr Val Thr Gln Asn Glu

- Asp Ala Asp Arg Asn Asn Ser Ser Arg Ile Asn Leu Gln Asp Ile Cys 200 Lys Asp Leu Asn Gln Glu Gly Asp Asp Gln Met Phe Val Arg Lys Thr 215 Ser Arg Glu Ile Gln Gly Gln Asn Arg Arg Asn Leu Ile His Glu Asp 235 230 Ile Ala Lys Ala Ile Lys Gln Leu Trp Asn Asn Asp Lys Gly Ile Lys 250 Gln Cys Phe Ala Arg Ser Asn Glu Phe Gln Leu Glu Gly Ser Ala Ala 265 260 Tyr Tyr Phe Asp Asn Ile Glu Lys Phe Ala Ser Pro Asn Tyr Val Cys Thr Asp Glu Asp Ile Leu Lys Gly Arg Ile Lys Thr Thr Gly Ile Thr 290 Glu Thr Glu Phe Asn Ile Gly Ser Ser Lys Phe Lys Val Leu Asp Ala 315 Gly Gln Arg Ser Glu Arg Lys Lys Trp Ile His Cys Phe Glu Gly 330 Ile Thr Ala Val Leu Phe Val Leu Ala Met Ser Glu Tyr Asp Gln Met 340 345 Leu Phe Glu Asp Glu Arg Val Asn Arg Met His Glu Ser Ile Met Leu 360 Phe Asp Thr Leu Leu Asn Ser Lys Trp Phe Lys Asp Thr Pro Phe Ile 375 Leu Phe Leu Asn Lys Ile Asp Leu Phe Glu Glu Lys Val Lys Ser Met 395 390
- Pro Ile Arg Lys Tyr Phe Pro Asp Tyr Gln Gly Arg Val Gly Asp Ala 405 410 415
- Glu Ala Gly Leu Lys Tyr Phe Glu Lys Ile Phe Leu Ser Leu Asn Lys 420 425 430
- Thr Asn Lys Pro Ile Tyr Val Lys Arg Thr Cys Ala Thr Asp Thr Gln 435 440 445
- Thr Met Lys Phe Val Leu Ser Ala Val Thr Asp Leu Ile Ile Gln Gln 450 455 460

Asn Leu Lys Gln Tyr Glu Leu Leu 465 470 <211> 472

<212> PRT

<213> Chimaera sp.

<400> 111

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro 1 5 10 15

Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln
20 25 30

Leu Glu Lys Gln Arg Asp Lys Asn Glu Ile Lys Leu Leu Leu Gly 35 40 45

Ala Gly Glu Ser Gly Lys Ser Thr Val Leu Lys Gln Leu Lys Leu Leu 50 55 60

His Gln Gly Gly Phe Ser His Gln Glu Arg Leu Gln Tyr Ala Gln Val 65 70 75 80

Ile Trp Ala Asp Ala Ile Gln Ser Met Lys Ile Leu Ile Ile Gln Ala 85 90 95

Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp Asp Pro Ile Asn Asn Lys
100 105 110

Asp Leu Phe Ala Cys Lys Arg Ile Leu Leu Lys Ala Lys Ala Leu Asp 115 120 125

Tyr Ile Asn Ala Ser Val Ala Gly Gly Ser Asp Phe Leu Asn Asp Tyr 130 135 140

Val Leu Lys Tyr Ser Glu Arg Tyr Glu Thr Arg Arg Arg Val Gln Ser 145 150 155 160

Thr Gly Arg Ala Lys Ala Ala Phe Asp Glu Asp Gly Asn Ile Ser Asn 165 170 175

Val Lys Ser Asp Thr Asp Arg Asp Ala Glu Thr Val Thr Gln Asn Glu 180 185 190

Asp Ala Asp Arg Asn Asn Ser Ser Arg Ile Asn Leu Gln Asp Ile Cys 195 200 205

Lys Asp Leu Asn Gln Glu Gly Asp Asp Gln Met Phe Val Arg Lys Thr 210 215 220

Ser Arg Glu Ile Gln Gly Gln Asn Arg Arg Asn Leu Ile His Glu Asp 225 230 235 240

Ile Ala Lys Ala Ile Lys Gln Leu Trp Asn Asn Asp Lys Gly Ile Lys
245 250 255

Gln Cys Phe Ala Arg Ser Asn Glu Phe Gln Leu Glu Gly Ser Ala Ala 260 265 270

Tyr Tyr Phe Asp Asn Ile Glu Lys Phe Ala Ser Pro Asn Tyr Val Cys 275 280 285 Thr Asp Glu Asp Ile Leu Lys Gly Arg Ile Lys Thr Thr Gly Ile Thr 290 295 300

Glu Thr Glu Phe Asn Ile Gly Ser Ser Lys Phe Lys Val Leu Asp Ala 305 310 315 320

Gly Gly Gln Arg Ser Glu Arg Lys Lys Trp Ile His Cys Phe Glu Gly 325 330 335

Ile Thr Ala Val Leu Phe Val Leu Ala Met Ser Glu Tyr Asp Gln Met 340 345 350

Leu Phe Glu Asp Glu Arg Val Asn Arg Met His Glu Ser Ile Met Leu 355 360 365

Phe Asp Thr Leu Leu Asn Ser Lys Trp Phe Lys Asp Thr Pro Phe Ile 370 375 380

Leu Phe Leu Asn Lys Ile Asp Leu Phe Glu Glu Lys Val Lys Ser Met 385 390 395 400

Pro Ile Arg Lys Tyr Phe Pro Asp Tyr Gln Gly Arg Val Gly Asp Ala 405 410 415

Glu Ala Gly Leu Lys Tyr Phe Glu Lys Ile Phe Leu Ser Leu Asn Lys 420 425 430

Thr Asn Lys Pro Ile Tyr Val Lys Arg Thr Cys Ala Thr Asp Thr Gln
435 440 445

Thr Met Lys Phe Val Leu Ser Ala Val Thr Asp Leu Ile Ile Gln Gln 450 455 460

Asn Leu Lys Gln Leu Met Leu Gln 465 470

<210> 112

<211> 472

<212> PRT

<213> Chimaera sp.

<400> 112

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro 1 5 10 15

Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln 20 25 30

Leu Glu Lys Gln Arg Asp Lys Asn Glu Ile Lys Leu Leu Leu Gly
35 40 45

Ala Gly Glu Ser Gly Lys Ser Thr Val Leu Lys Gln Leu Lys Leu Leu 50 55 60

His Gln Gly Gly Phe Ser His Gln Glu Arg Leu Gln Tyr Ala Gln Val

- Ile Trp Ala Asp Ala Ile Gln Ser Met Lys Ile Leu Ile Ile Gln Ala 85 90 95
- Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp Asp Pro Ile Asn Asn Lys
  100 105 110
- Asp Leu Phe Ala Cys Lys Arg Ile Leu Leu Lys Ala Lys Ala Leu Asp
- Tyr Ile Asn Ala Ser Val Ala Gly Gly Ser Asp Phe Leu Asn Asp Tyr 130 135 140
- Val Leu Lys Tyr Ser Glu Arg Tyr Glu Thr Arg Arg Arg Val Gln Ser 145 150 155 160
- Thr Gly Arg Ala Lys Ala Ala Phe Asp Glu Asp Gly Asn Ile Ser Asn 165 170 175
- Val Lys Ser Asp Thr Asp Arg Asp Ala Glu Thr Val Thr Gln Asn Glu 180 185 190
- Asp Ala Asp Arg Asn Asn Ser Ser Arg Ile Asn Leu Gln Asp Ile Cys 195 200 205
- Lys Asp Leu Asn Gln Glu Gly Asp Asp Gln Met Phe Val Arg Lys Thr 210 215 220
- Ser Arg Glu Ile Gln Gly Gln Asn Arg Arg Asn Leu Ile His Glu Asp 225 230 235 240
- Ile Ala Lys Ala Ile Lys Gln Leu Trp Asn Asn Asp Lys Gly Ile Lys 245 250 255
- Gln Cys Phe Ala Arg Ser Asn Glu Phe Gln Leu Glu Gly Ser Ala Ala 260 265 270
- Tyr Tyr Phe Asp Asn Ile Glu Lys Phe Ala Ser Pro Asn Tyr Val Cys 275 280 285
- Thr Asp Glu Asp Ile Leu Lys Gly Arg Ile Lys Thr Thr Gly Ile Thr 290 295 300
- Glu Thr Glu Phe Asn Ile Gly Ser Ser Lys Phe Lys Val Leu Asp Ala 305 310 315 320
- Gly Gln Arg Ser Glu Arg Lys Lys Trp Ile His Cys Phe Glu Gly 325 330 335
- Ile Thr Ala Val Leu Phe Val Leu Ala Met Ser Glu Tyr Asp Gln Met 340 345 350
- Leu Phe Glu Asp Glu Arg Val Asn Arg Met His Glu Ser Ile Met Leu 355 360 365
- Phe Asp Thr Leu Leu Asn Ser Lys Trp Phe Lys Asp Thr Pro Phe Ile 370 375 380

Leu Phe Leu Asn Lys Ile Asp Leu Phe Glu Glu Lys Val Lys Ser Met 385 390 395 400

Pro Ile Arg Lys Tyr Phe Pro Asp Tyr Gln Gly Arg Val Gly Asp Ala 405 410 415

Glu Ala Gly Leu Lys Tyr Phe Glu Lys Ile Phe Leu Ser Leu Asn Lys 420 425 430

Thr Asn Lys Pro Ile Tyr Val Lys Arg Thr Cys Ala Thr Asp Thr Gln 435 440 445

Thr Met Lys Phe Val Leu Ser Ala Val Thr Asp Leu Ile Ile Gln Gln 450 455 460

Asn Leu Lys Tyr Ile Gly Leu Cys 465 470

<210> 113

1

<211> 472

<212> PRT

<213> Chimaera sp.

<400> 113

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro 1 5 10 15

Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln 20 25 30

Leu Glu Lys Gln Arg Asp Lys Asn Glu Ile Lys Leu Leu Leu Gly 35 40 45

Ala Gly Glu Ser Gly Lys Ser Thr Val Leu Lys Gln Leu Lys Leu Leu 50 55 60

His Gln Gly Gly Phe Ser His Gln Glu Arg Leu Gln Tyr Ala Gln Val
65 70 75 80

Ile Trp Ala Asp Ala Ile Gln Ser Met Lys Ile Leu Ile Ile Gln Ala 85 90 95

Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp Asp Pro Ile Asn Asn Lys 100 105 110

Asp Leu Phe Ala Cys Lys Arg Ile Leu Leu Lys Ala Lys Ala Leu Asp 115 120 125

Tyr Ile Asn Ala Ser Val Ala Gly Gly Ser Asp Phe Leu Asn Asp Tyr 130 135 140

Val Leu Lys Tyr Ser Glu Arg Tyr Glu Thr Arg Arg Arg Val Gln Ser 145 150 155 160

Thr Gly Arg Ala Lys Ala Ala Phe Asp Glu Asp Gly Asn Ile Ser Asn 165 170 175

- Val Lys Ser Asp Thr Asp Arg Asp Ala Glu Thr Val Thr Gln Asn Glu 180 185 190
- Asp Ala Asp Arg Asn Asn Ser Ser Arg Ile Asn Leu Gln Asp Ile Cys 195 200 205
- Lys Asp Leu Asn Gln Glu Gly Asp Asp Gln Met Phe Val Arg Lys Thr 210 215 220
- Ser Arg Glu Ile Gln Gly Gln Asn Arg Arg Asn Leu Ile His Glu Asp 225 230 235 240
- Ile Ala Lys Ala Ile Lys Gln Leu Trp Asn Asn Asp Lys Gly Ile Lys 245 250 255
- Gln Cys Phe Ala Arg Ser Asn Glu Phe Gln Leu Glu Gly Ser Ala Ala 260 265 270
- Tyr Tyr Phe Asp Asn Ile Glu Lys Phe Ala Ser Pro Asn Tyr Val Cys 275 280 285
- Thr Asp Glu Asp Ile Leu Lys Gly Arg Ile Lys Thr Thr Gly Ile Thr 290 295 300
- Glu Thr Glu Phe Asn Ile Gly Ser Ser Lys Phe Lys Val Leu Asp Ala 305 310 315 320
- Gly Gln Arg Ser Glu Arg Lys Lys Trp Ile His Cys Phe Glu Gly 325 330 335
- Ile Thr Ala Val Leu Phe Val Leu Ala Met Ser Glu Tyr Asp Gln Met 340 345 350
- Leu Phe Glu Asp Glu Arg Val Asn Arg Met His Glu Ser Ile Met Leu 355 360 365
- Phe Asp Thr Leu Leu Asn Ser Lys Trp Phe Lys Asp Thr Pro Phe Ile 370 375 380
- Leu Phe Leu Asn Lys Ile Asp Leu Phe Glu Glu Lys Val Lys Ser Met 385 390 395 400
- Pro Ile Arg Lys Tyr Phe Pro Asp Tyr Gln Gly Arg Val Gly Asp Ala 405 410 415
- Glu Ala Gly Leu Lys Tyr Phe Glu Lys Ile Phe Leu Ser Leu Asn Lys 420 425 430
- Thr Asn Lys Pro Ile Tyr Val Lys Arg Thr Cys Ala Thr Asp Thr Gln 435 440 445
- Thr Met Lys Phe Val Leu Ser Ala Val Thr Asp Leu Ile Ile Gln Gln 450 455 460

Asn Leu Lys Gly Cys Gly Leu Tyr 465 470

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<210> 114
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<211> 472

<212> PRT

<213> Chimaera sp.

## <400> 114

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro 1 5 10 15

Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln 20 25 30

Leu Glu Lys Gln Arg Asp Lys Asn Glu Ile Lys Leu Leu Leu Gly
35 40 45

Ala Gly Glu Ser Gly Lys Ser Thr Val Leu Lys Gln Leu Lys Leu Leu 50 55 60

His Gln Gly Gly Phe Ser His Gln Glu Arg Leu Gln Tyr Ala Gln Val 65 70 75 80

Ile Trp Ala Asp Ala Ile Gln Ser Met Lys Ile Leu Ile Ile Gln Ala 85 90 95

Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp Asp Pro Ile Asn Asn Lys
100 105 110

Asp Leu Phe Ala Cys Lys Arg Ile Leu Leu Lys Ala Lys Ala Leu Asp 115 120 125

Tyr Ile Asn Ala Ser Val Ala Gly Gly Ser Asp Phe Leu Asn Asp Tyr 130 135 140

Val Leu Lys Tyr Ser Glu Arg Tyr Glu Thr Arg Arg Arg Val Gln Ser 145 150 155 160

Thr Gly Arg Ala Lys Ala Ala Phe Asp Glu Asp Gly Asn Ile Ser Asn 165 170 175

Val Lys Ser Asp Thr Asp Arg Asp Ala Glu Thr Val Thr Gln Asn Glu 180 185 190

Asp Ala Asp Arg Asn Asn Ser Ser Arg Ile Asn Leu Gln Asp Ile Cys 195 200 205

Lys Asp Leu Asn Gln Glu Gly Asp Asp Gln Met Phe Val Arg Lys Thr 210 215 220

Ser Arg Glu Ile Gln Gly Gln Asn Arg Arg Asn Leu Ile His Glu Asp 225 230 235 240

Ile Ala Lys Ala Ile Lys Gln Leu Trp Asn Asn Asp Lys Gly Ile Lys 245 250 255

Gln Cys Phe Ala Arg Ser Asn Glu Phe Gln Leu Glu Gly Ser Ala Ala 260 265 270

Tyr Tyr Phe Asp Asn Ile Glu Lys Phe Ala Ser Pro Asn Tyr Val Cys

Thr Asp Glu Asp Ile Leu Lys Gly Arg Ile Lys Thr Thr Gly Ile Thr 290 295 300

Glu Thr Glu Phe Asn Ile Gly Ser Ser Lys Phe Lys Val Leu Asp Ala 305 310 315 320

Gly Gly Gln Arg Ser Glu Arg Lys Lys Trp Ile His Cys Phe Glu Gly
325 330 335

Ile Thr Ala Val Leu Phe Val Leu Ala Met Ser Glu Tyr Asp Gln Met 340 345 350

Leu Phe Glu Asp Glu Arg Val Asn Arg Met His Glu Ser Ile Met Leu 355 360 . 365

Phe Asp Thr Leu Leu Asn Ser Lys Trp Phe Lys Asp Thr Pro Phe Ile 370 375 380

Leu Phe Leu Asn Lys Ile Asp Leu Phe Glu Glu Lys Val Lys Ser Met 385 390 395 400

Pro Ile Arg Lys Tyr Phe Pro Asp Tyr Gln Gly Arg Val Gly Asp Ala
405 410 415

Glu Ala Gly Leu Lys Tyr Phe Glu Lys Ile Phe Leu Ser Leu Asn Lys 420 425 430

Thr Asn Lys Pro Ile Tyr Val Lys Arg Thr Cys Ala Thr Asp Thr Gln
435 440 445

Thr Met Lys Phe Val Leu Ser Ala Val Thr Asp Leu Ile Ile Gln Gln
450 455 460

Asn Leu Asp Glu Ile Asn Leu Leu 465 470

<210> 115

<211> 472

<212> PRT

<213> Chimaera sp.

<400> 115

Met Gly Cys Thr Val Ser Thr Gln Thr Ile Gly Asp Glu Ser Asp Pro 1 5 10 15

Phe Leu Gln Asn Lys Arg Ala Asn Asp Val Ile Glu Gln Ser Leu Gln 20 25 30

Leu Glu Lys Gln Arg Asp Lys Asn Glu Ile Lys Leu Leu Leu Gly
35 40 45

Ala Gly Glu Ser Gly Lys Ser Thr Val Leu Lys Gln Leu Lys Leu Leu 50 55 60

His Gln Gly Gly Phe Ser His Gln Glu Arg Leu Gln Tyr Ala Gln Val

Ile Trp Ala Asp Ala Ile Gln Ser Met Lys Ile Leu Ile Ile Gln Ala 85 90 95

65

Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp Asp Pro Ile Asn Asn Lys
100 105 110

Asp Leu Phe Ala Cys Lys Arg Ile Leu Leu Lys Ala Lys Ala Leu Asp 115 120 125

Tyr Ile Asn Ala Ser Val Ala Gly Gly Ser Asp Phe Leu Asn Asp Tyr 130 135 140

Val Leu Lys Tyr Ser Glu Arg Tyr Glu Thr Arg Arg Arg Val Gln Ser 145 150 155 160

Thr Gly Arg Ala Lys Ala Ala Phe Asp Glu Asp Gly Asn Ile Ser Asn 165 170 175

Val Lys Ser Asp Thr Asp Arg Asp Ala Glu Thr Val Thr Gln Asn Glu 180 185 190

Asp Ala Asp Arg Asn Asn Ser Ser Arg Ile Asn Leu Gln Asp Ile Cys 195 200 205

Lys Asp Leu Asn Gln Glu Gly Asp Asp Gln Met Phe Val Arg Lys Thr 210 215 220

Ser Arg Glu Ile Gln Gly Gln Asn Arg Arg Asn Leu Ile His Glu Asp 225 230 235 240

Ile Ala Lys Ala Ile Lys Gln Leu Trp Asn Asn Asp Lys Gly Ile Lys 245 250 255

Gln Cys Phe Ala Arg Ser Asn Glu Phe Gln Leu Glu Gly Ser Ala Ala 260 265 270

Tyr Tyr Phe Asp Asn Ile Glu Lys Phe Ala Ser Pro Asn Tyr Val Cys 275 280 285

Thr Asp Glu Asp Ile Leu Lys Gly Arg Ile Lys Thr Thr Gly Ile Thr 290 295 300

Glu Thr Glu Phe Asn Ile Gly Ser Ser Lys Phe Lys Val Leu Asp Ala 305 310 315 320

Gly Gly Gln Arg Ser Glu Arg Lys Lys Trp Ile His Cys Phe Glu Gly 325 330 335

Ile Thr Ala Val Leu Phe Val Leu Ala Met Ser Glu Tyr Asp Gln Met 340 345 350

Leu Phe Glu Asp Glu Arg Val Asn Arg Met His Glu Ser Ile Met Leu 355 360 365

Phe Asp Thr Leu Leu Asn Ser Lys Trp Phe Lys Asp Thr Pro Phe Ile 370 375 380

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Leu Phe Leu Asn Lys Ile Asp Leu Phe Glu Glu Lys Val Lys Ser Met
                    390
Pro Ile Arg Lys Tyr Phe Pro Asp Tyr Gln Gly Arg Val Gly Asp Ala
                405
Glu Ala Gly Leu Lys Tyr Phe Glu Lys Ile Phe Leu Ser Leu Asn Lys
                                 425
            420
Thr Asn Lys Pro Ile Tyr Val Lys Arg Thr Cys Ala Thr Asp Thr Gln
Thr Met Lys Phe Val Leu Ser Ala Val Thr Asp Leu Ile Ile Gln Gln
                                             460
                        455
Asn Leu Arg Gln Tyr Glu Leu Leu
                    470
<210> 116
<211> 67
<212> DNA
<213> Chimaera sp.
<400> 116
acgtggtctc ccatgacttt ggaatctatt atggcttgtt gtcttagtac gcaaacaata 60
ggagacg
<210> 117
<211> 21
<212> DNA
<213> Chimaera sp.
<400> 117
                                                                    21
gtatctttga accacttaga g
<210> 118
<211> 478
<212> PRT
<213> Chimaera sp.
<400> 118
Met Thr Leu Glu Ser Ile Met Ala Cys Cys Leu Ser Thr Gln Thr Ile
                   5
                                      10
Gly Asp Glu Ser Asp Pro Phe Leu Gln Asn Lys Arg Ala Asn Asp Val
Ile Glu Gln Ser Leu Gln Leu Glu Lys Gln Arg Asp Lys Asn Glu Ile
                              40
Lys Leu Leu Leu Gly Ala Gly Glu Ser Gly Lys Ser Thr Val Leu
      50
Lys Gln Leu Lys Leu Leu His Gln Gly Gly Phe Ser His Gln Glu Arg
                      70
                                           75
 65
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Leu Gln Tyr Ala Gln Val Ile Trp Ala Asp Ala Ile Gln Ser Met Lys 85 90 95

2

Ile Leu Ile Ile Gln Ala Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp 100 105 110

Asp Pro Ile Asn Asn Lys Asp Leu Phe Ala Cys Lys Arg Ile Leu Leu 115 120 125

Lys Ala Lys Ala Leu Asp Tyr Ile Asn Ala Ser Val Ala Gly Gly Ser 130 135 140

Asp Phe Leu Asn Asp Tyr Val Leu Lys Tyr Ser Glu Arg Tyr Glu Thr 145 150 155 160

Arg Arg Arg Val Gln Ser Thr Gly Arg Ala Lys Ala Ala Phe Asp Glu 165 170 175

Asp Gly Asn Ile Ser Asn Val Lys Ser Asp Thr Asp Arg Asp Ala Glu 180 185 190

Thr Val Thr Gln Asn Glu Asp Ala Asp Arg Asn Asn Ser Ser Arg Ile 195 200 205

Asn Leu Gln Asp Ile Cys Lys Asp Leu Asn Gln Glu Gly Asp Asp Gln 210 215 220

Met Phe Val Arg Lys Thr Ser Arg Glu Ile Gln Gly Gln Asn Arg Arg 225 230 235 240

Asn Leu Ile His Glu Asp Ile Ala Lys Ala Ile Lys Gln Leu Trp Asn 245 250 255

Asn Asp Lys Gly Ile Lys Gln Cys Phe Ala Arg Ser Asn Glu Phe Gln 260 265 270

Leu Glu Gly Ser Ala Ala Tyr Tyr Phe Asp Asn Ile Glu Lys Phe Ala 275 280 285

Ser Pro Asn Tyr Val Cys Thr Asp Glu Asp Ile Leu Lys Gly Arg Ile 290 295 300

Lys Thr Thr Gly Ile Thr Glu Thr Glu Phe Asn Ile Gly Ser Ser Lys 305 310 315 320

Phe Lys Val Leu Asp Ala Gly Gly Gln Arg Ser Glu Arg Lys Lys Trp 325 330 335

Ile His Cys Phe Glu Gly Ile Thr Ala Val Leu Phe Val Leu Ala Met 340 345 350

Ser Glu Tyr Asp Gln Met Leu Phe Glu Asp Glu Arg Val Asn Arg Met 355 360 365

His Glu Ser Ile Met Leu Phe Asp Thr Leu Leu Asn Ser Lys Trp Phe 370 375 380

Lys Asp Thr Pro Phe Ile Leu Phe Leu Asn Lys Ile Asp Leu Phe Glu

Met Thr Leu Glu Ser Ile Met Ala Cys Cys Leu Ser Glu Glu Ala Lys

1

Glu Ala Arg Arg Ile Asn Asp Val Ile Glu Gln Ser Leu Gln Leu Glu 25 20 Lys Gln Arg Asp Lys Asn Glu Ile Lys Leu Leu Leu Gly Ala Gly 40 Glu Ser Gly Lys Ser Thr Val Leu Lys Gln Leu Lys Leu Leu His Gln Gly Gly Phe Ser His Gln Glu Arg Leu Gln Tyr Ala Gln Val Ile Trp Ala Asp Ala Ile Gln Ser Met Lys Ile Leu Ile Ile Gln Ala Arg Lys Leu Gly Ile Gln Leu Asp Cys Asp Asp Pro Ile Asn Asn Lys Asp Leu 105 Phe Ala Cys Lys Arg Ile Leu Leu Lys Ala Lys Ala Leu Asp Tyr Ile 120 Asn Ala Ser Val Ala Gly Gly Ser Asp Phe Leu Asn Asp Tyr Val Leu 135 Lys Tyr Ser Glu Arg Tyr Glu Thr Arg Arg Arg Val Gln Ser Thr Gly 150 145 Arg Ala Lys Ala Ala Phe Asp Glu Asp Gly Asn Ile Ser Asn Val Lys 170 Ser Asp Thr Asp Arg Asp Ala Glu Thr Val Thr Gln Asn Glu Asp Ala 185 Asp Arg Asn Asn Ser Ser Arg Ile Asn Leu Gln Asp Ile Cys Lys Asp 205 195 200 Leu Asn Gln Glu Gly Asp Asp Gln Met Phe Val Arg Lys Thr Ser Arg 215 Glu Ile Gln Gly Gln Asn Arg Arg Asn Leu Ile His Glu Asp Ile Ala 225 Lys Ala Ile Lys Gln Leu Trp Asn Asn Asp Lys Gly Ile Lys Gln Cys 250

Phe Ala Arg Ser Asn Glu Phe Gln Leu Glu Gly Ser Ala Ala Tyr Tyr 260 265 270

Phe Asp Asn Ile Glu Lys Phe Ala Ser Pro Asn Tyr Val Cys Thr Asp 275 280 285

Glu Asp Ile Leu Lys Gly Arg Ile Lys Thr Thr Gly Ile Thr Glu Thr 290 295 300

Glu Phe Asn Ile Gly Ser Ser Lys Phe Lys Val Leu Asp Ala Gly Gly 305 310 315 320

- Gln Arg Ser Glu Arg Lys Lys Trp Ile His Cys Phe Glu Gly Ile Thr 325 330 335
- Ala Val Leu Phe Val Leu Ala Met Ser Glu Tyr Asp Gln Met Leu Phe 340 345 350
- Glu Asp Glu Arg Val Asn Arg Met His Glu Ser Ile Met Leu Phe Asp 355 360 365
- Thr Leu Leu Asn Ser Lys Trp Phe Lys Asp Thr Pro Phe Ile Leu Phe 370 375 380
- Leu Asn Lys Ile Asp Leu Phe Glu Glu Lys Val Lys Ser Met Pro Ile 385 390 395 400
- Arg Lys Tyr Phe Pro Asp Tyr Gln Gly Arg Val Gly Asp Ala Glu Ala 405 410 415
- Gly Leu Lys Tyr Phe Glu Lys Ile Phe Leu Ser Leu Asn Lys Thr Asn 420 425 430
- Lys Pro Ile Tyr Val Lys Arg Thr Cys Ala Thr Asp Thr Gln Thr Met 435 440 445
- Lys Phe Val Leu Ser Ala Val Thr Asp Leu Ile Ile Gln Gln Asn Leu 450 455 460

Lys Glu Tyr Asn Leu Val 465 470